



att ccg gct tct atg gag cac tcg gga cca ggt ccg cgg cgc gcg cac tcg ctc
gct cgc cgc ccc cca gcc agc tct cgc ttc cgc gcc gac cgc gcc ccc cgc
ctc ctc gct gca ccc cgc gac cta gag cca aga aag ttt gtg tgg cga gtg agg
gcc gga gag gag agc gcg ccc gcg gag tgc cgt cca gac cag cgc ggc ccc ggc
gga gag ggg agc gcc ccg agc cca ggc ggc ggc tag ccc gag tcc gcg acc
-26 -20
Met Gly Ala Ala Ala Arg Ser Leu Pro Leu Ala Phe
ccc gcc cct ccg ccc gcc atg ggcc gcc gcc cgcc agc ctg ccg ctc gcg ttc
-10 -1 1
Cys Leu Leu Leu Leu Gly Thr Leu Leu Pro Arg Ala Asp Ala Cys Ser Cys Ser
tgc ctc ctg ctg ggg acg ctg ctc ccc cgg gcc gac gcc tgc agc tgc tcc
10 20
Pro Val His Pro Gln Gln Ala Phe Cys Asn Ala Asp Ile Val Ile Arg Ala Lys
ccg gtg cac ccg caa cag gcg ttt tgc aat gca gac ata gtg atc agg gcc aaa
30 40
Ala Val Asn Lys Lys Glu Val Asp Ser Gly Asn Asp Ile Tyr Gly Asn Pro Ile
gca gtc aat aag aag gag gtg gac tct ggcc aac gac atc tac ggcc aac ccc atc
50
Lys Arg Ile Gln Tyr Glu Ile Lys Gln Ile Lys Met Phe Lys Gly Pro Asp Gln
aag cgg att cag tat gag atc aag cag ata aag atg ttc aag ggaa cct gat cag
60 70
Asp Ile Glu Phe Ile Tyr Thr Ala Pro Ala Ala Val Cys Gly Val Ser Leu
gac ata gag ttt atc tac aca gca ccc gcc gct gcc gtg tgt ggg gtc tgc ctg
80 90
Asp Ile Gly Gly Lys Lys Glu Tyr Leu Ile Ala Gly Lys Ala Glu Gly Asn Gly
gac att gga gga aag aag gag tat ctc att gca ggg aag gcc gag ggg aat ggc
100 110
Asn Met His Ile Thr Leu Cys Asp Phe Ile Val Pro Trp Asp Thr Leu Ser Ala
aat atg cat atc acc ctc tgt gac ttc atc gtg ccc tgg gac acc ctg agt gcc
120 130
Thr Gln Lys Lys Ser Leu Asn His Arg Tyr Gln Met Gly Cys Glu Cys Lys Ile
acc cag aag aag agc ctg aac cac agg tac cag atg ggc tgt gag tgc aag atc
140
Thr Arg Cys Pro Met Ile Pro Cys Tyr Ile Ser Ser Pro Asp Glu Cys Leu Trp
act cga tgc ccc atg atc cca tgc tac atc tcc tct ccg gac gag tgc ctc tgg
150 160
Met Asp Trp Val Thr Glu Lys Asn Ile Asn Gly His Gln Ala Lys Phe Phe Ala
atg gac tgg gtc acg gag aag aac atc aac gga cac cag ggc aag ttc ttc gcc
170 180
Cys Ile Lys Arg Ser Asp Gly Ser Cys Ala Trp Tyr Arg Gly Ala Ala Pro Pro
tgc atc aag aga agc gac ggc tcc tgc gcc tgg tac cgc gga gca gca ccc ccc
190 194
Lys Gln Glu Phe Leu Asp Ile Glu Asp Pro
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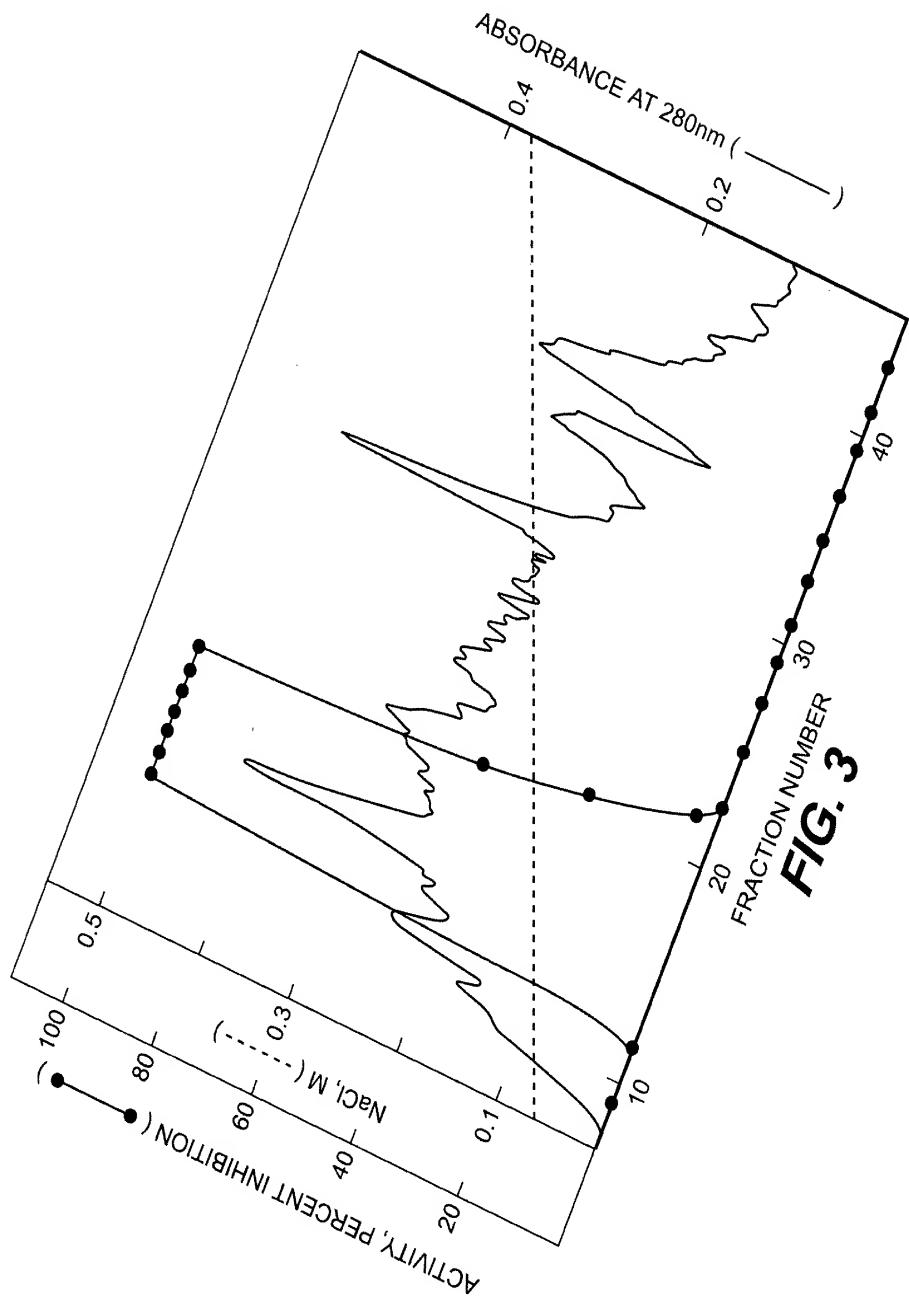
FIG. 1

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cag ccc ccc gc_g ccc cca gcc cgc cgc gag cag cgc ccc gac ccc cca
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gc_g gc_g gcc ccc gcc cgc cca gcc ccc cgg ccc gcc atg ggc gcc gc_g gcc ccc
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acc ctg cg_g ctg gc_g ctc ctg ctg ctg gc_g acg ctg ctt cgc ccc gcc
-1 1 10
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gac gcc tgc agc tcc ccc gtg cac ccc cag gc_g ttt tgc aat gca gat
20 30
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gta gtg atc agg gcc aaa gc_g gtc agt gag aag gaa gtg gac tct gg_a aac gac
40 50
Ile Tyr Gly Asn Pro Ile Lys Arg Ile Gln Tyr Glu Ile Lys Gln Ile Lys Met
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60 70
Phe Lys Gly Pro Glu Lys Asp Ile Glu Phe Ile Tyr Thr Ala Pro Ser Ser Ala
ttc aaa ggg cct gag aag gat ata gag ttt atc tac acg gcc ccc tcc tcg gca
80
Val Cys Gly Val Ser Leu Asp Val Gly Gly Lys Lys Glu Tyr Leu Ile Ala Gly
gtg tgt ggg gtc tcg ctg gac gtt gg_a gg_a aag aag gaa tat ctc att gca gg_a
90 100
Lys Ala Glu Gly Asp Gly Lys Met His Ile Thr Leu Cys Asp Phe Ile Val Pro
aag gcc gag ggg gac ggc aag atg cac atc acc ctc tgt gac ttc atc gtg ccc
110 120
Trp Asp Thr Leu Ser Thr Thr Gln Lys Lys Ser Leu Asn His Arg Tyr Gln Met
tgg gac acc ctg agc acc acc cag aag aag agc ctg aac cac agg tac cag atg
130 140
Gly Cys Glu Cys Lys Ile Thr Arg Cys Pro Met Ile Pro Cys Tyr Ile Ser Ser
ggc tgc gag tgc aag atc acg ccc tgc ccc atg atc ccg tgc tac atc tcc tcc
150 160
Pro Asp Glu Cys Leu Trp Met Asp Trp Val Thr Glu Lys Asn Ile Asn Gly His
ccg gac gag tgc ctc tgg atg gac tgg gtc aca gag aag aac atc aac ggg cac
170
Gln Ala Lys Phe Phe Ala Cys Ile Lys Arg Ser Asp Gly Ser Cys Ala Trp Tyr
cag gcc aag ttc ttc gcc tgc atc aag aag agt gac ggc tcc tgt ggc tgg tac
180 190 194
Arg Gly Ala Ala Pro Pro Lys Gln Glu Phe Leu Asp Ile Glu Asp Pro
cgc ggc gc_g ccc ccc aag cag gag ttt ctc gac atc gag gac cca taa gca
ggc ctc caa cgc ccc tgt ggc caa aaa aag cct cca agg gtt tcg act
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gga att c

FIG. 2



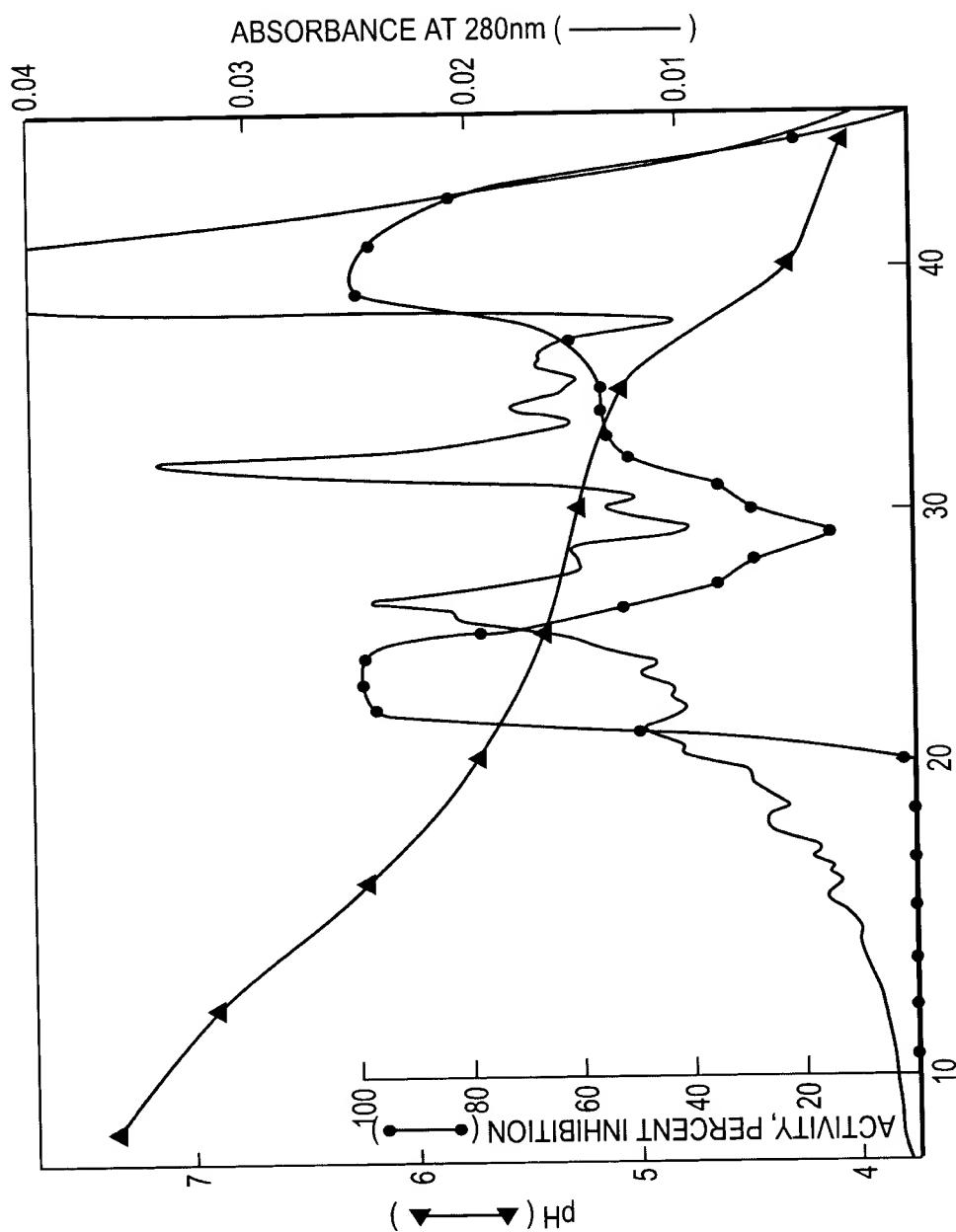
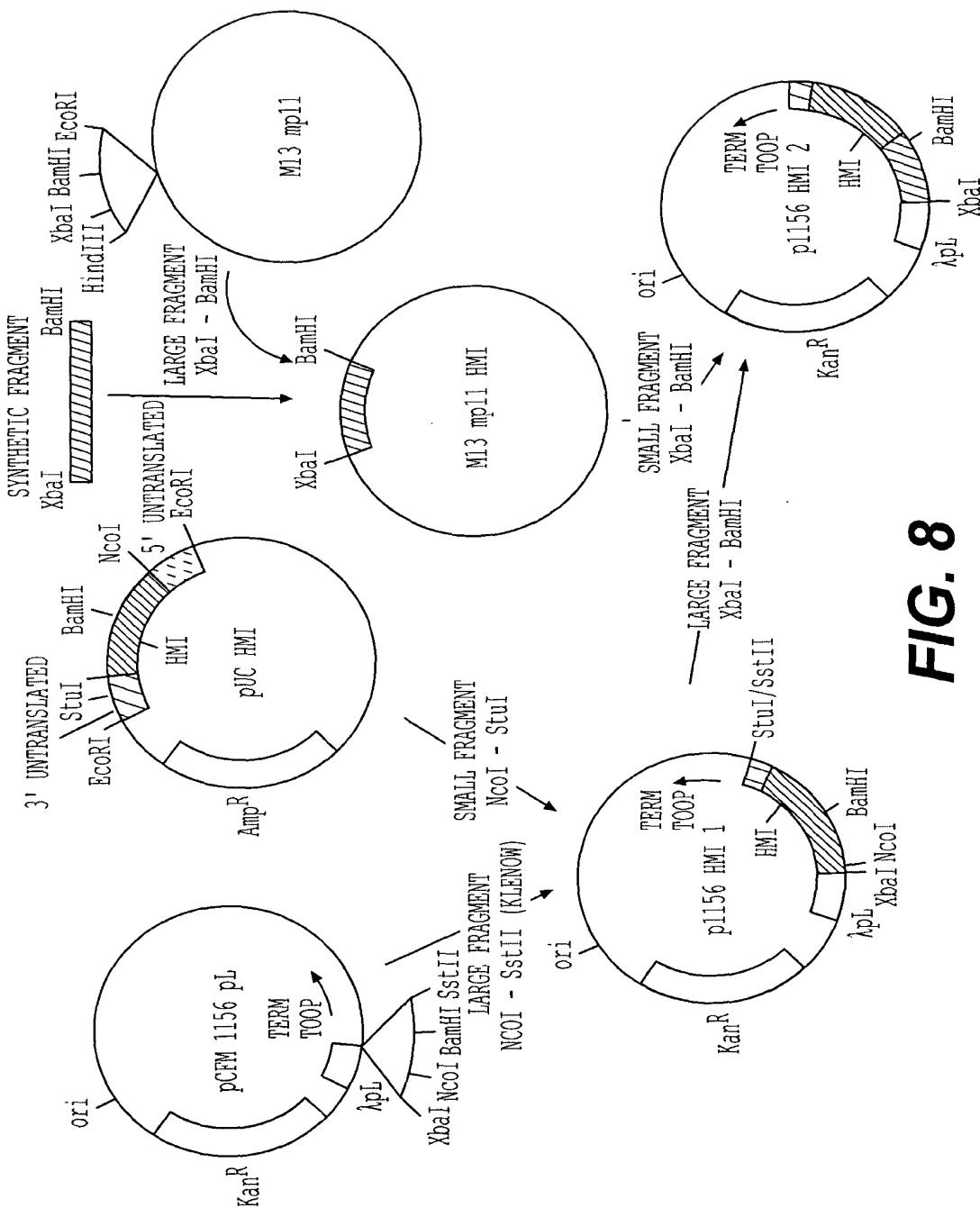


FIG. 4



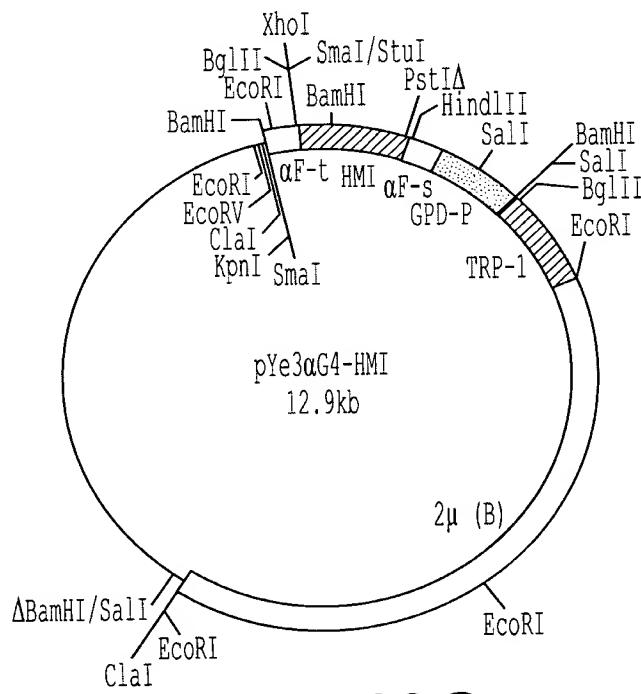
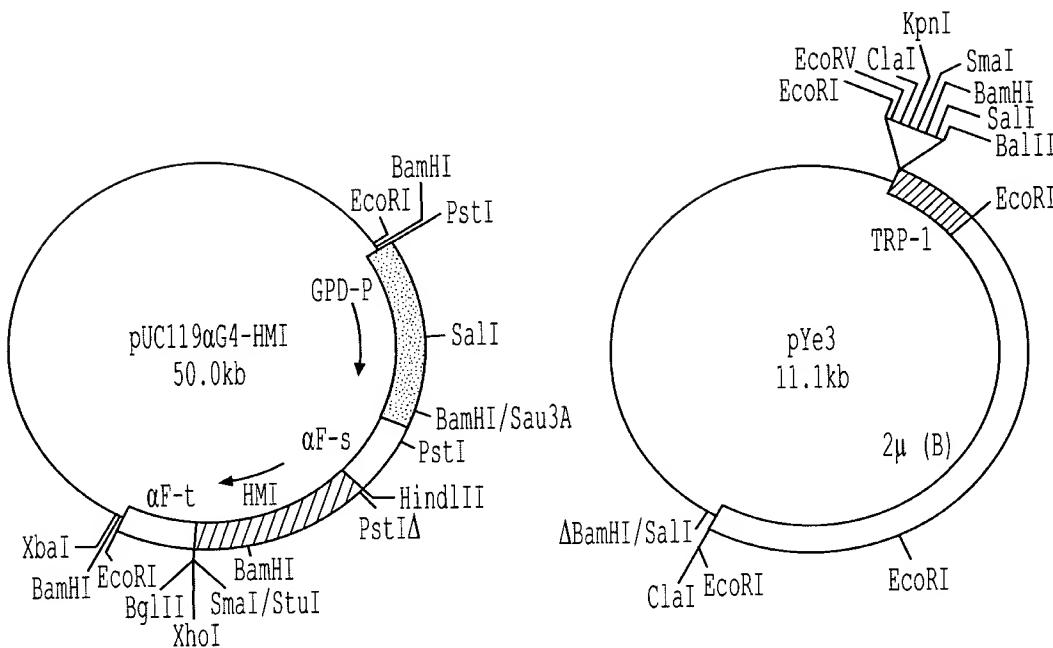


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TTTTTT GGTCCTCCA TTATTTATTA CACAAGAACAGAGGACATG TGGGAGTTGT

70 80 90 100 110 120
AGCTTTTGT AACGCTGATG TAGTTATCCG TGCAAAAGCT GTTTCTGAAA AAGAAGTTGA
TCGAAAAACA TTGCGACTAC ATCAATAGGC ACGTTTCGA CAAAGACTTT TTCTTCAACT

130 140 150 160
TTCTGGTAAC GACATCTACG GTAACCCGAT CAAAAG
AAGACCATTG CTGTAGATGC CATTGGGCTA GTTTCCCTAG

FIG. 9



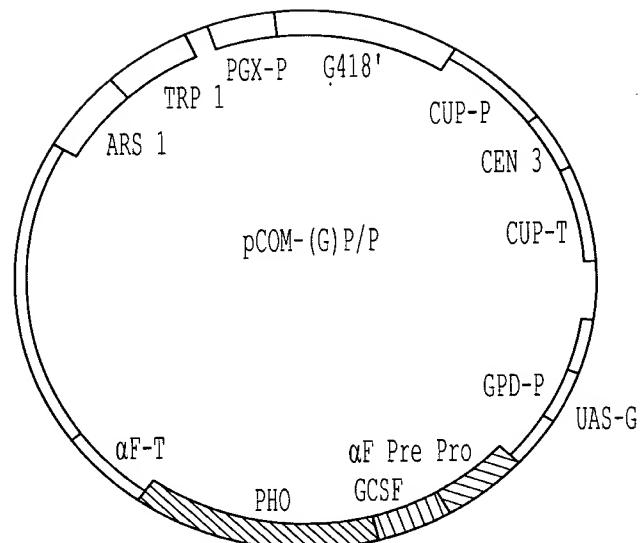


FIG. 11

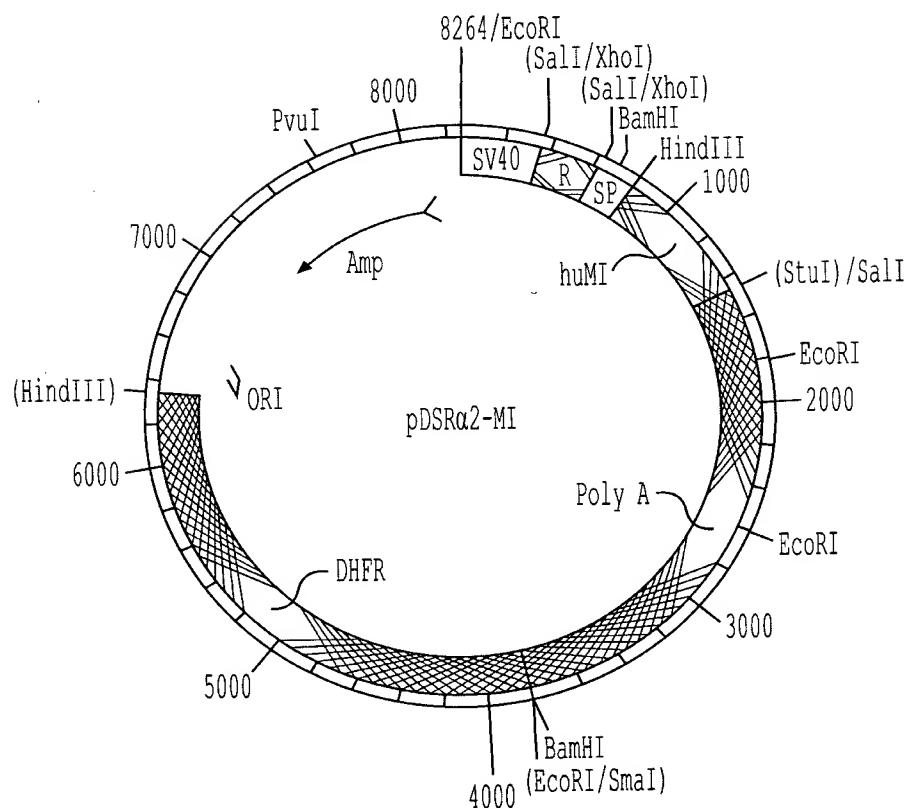


FIG. 12

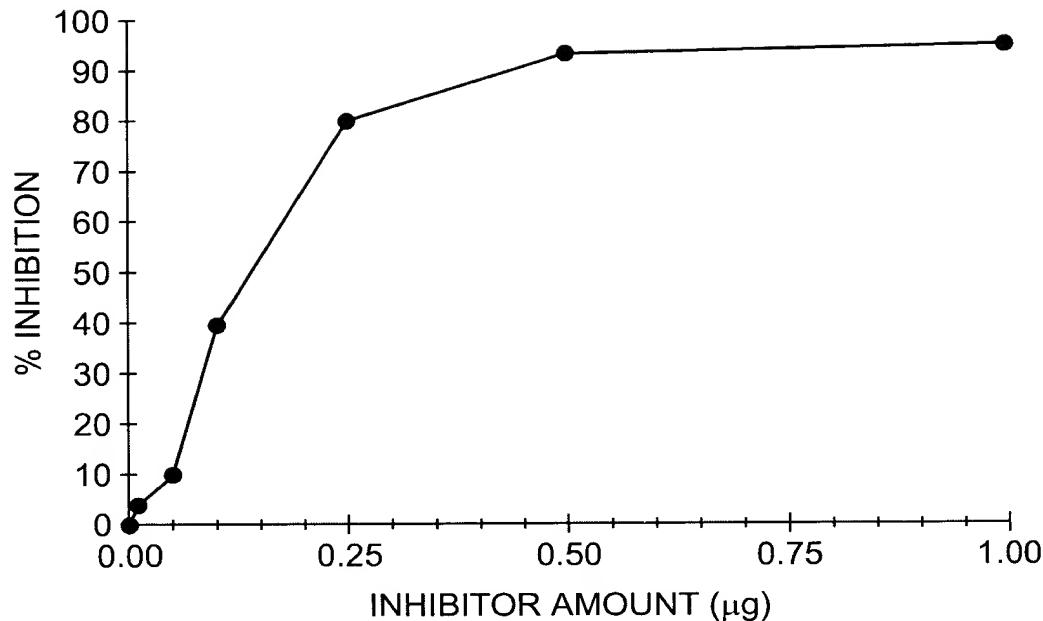


FIG. 15A

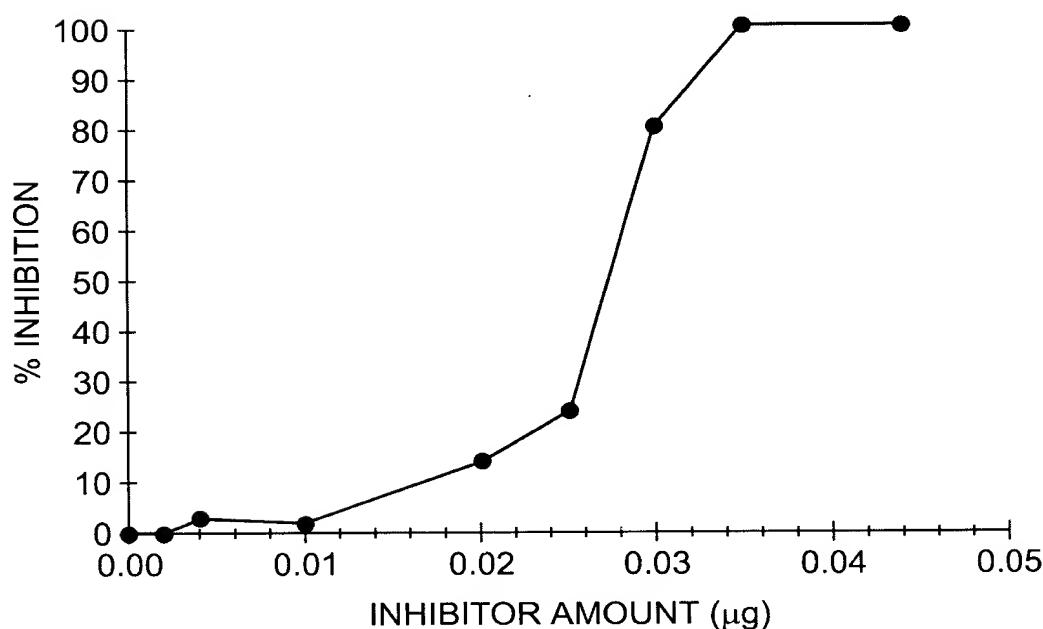


FIG. 15B

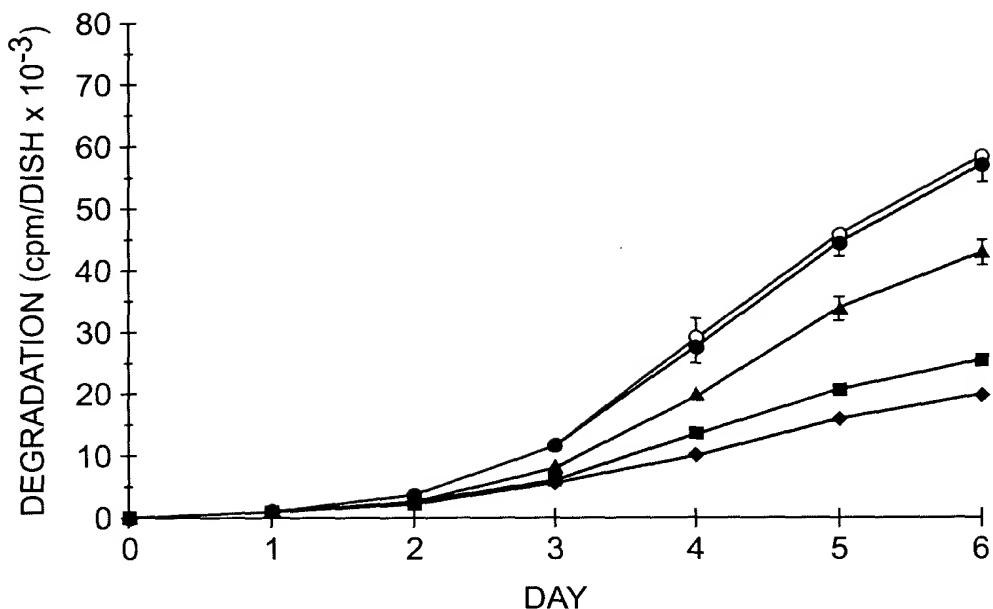


FIG. 16

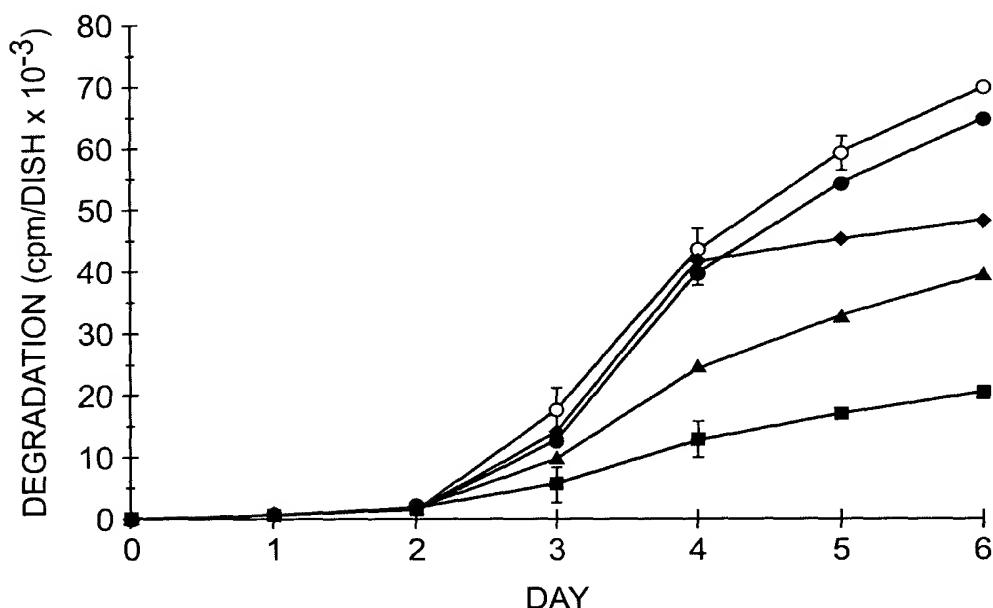


FIG. 17

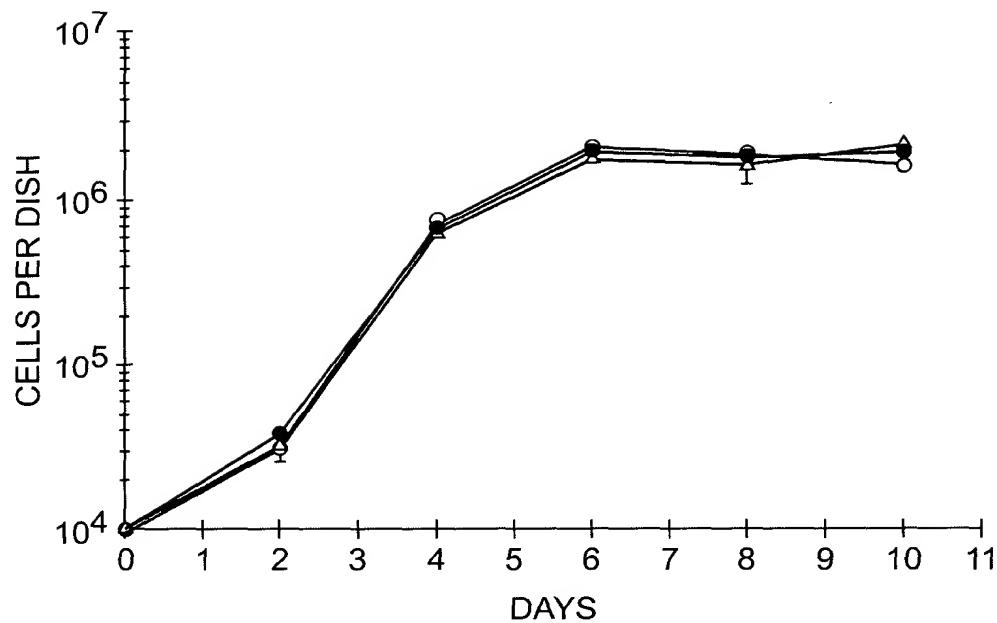


FIG. 18A

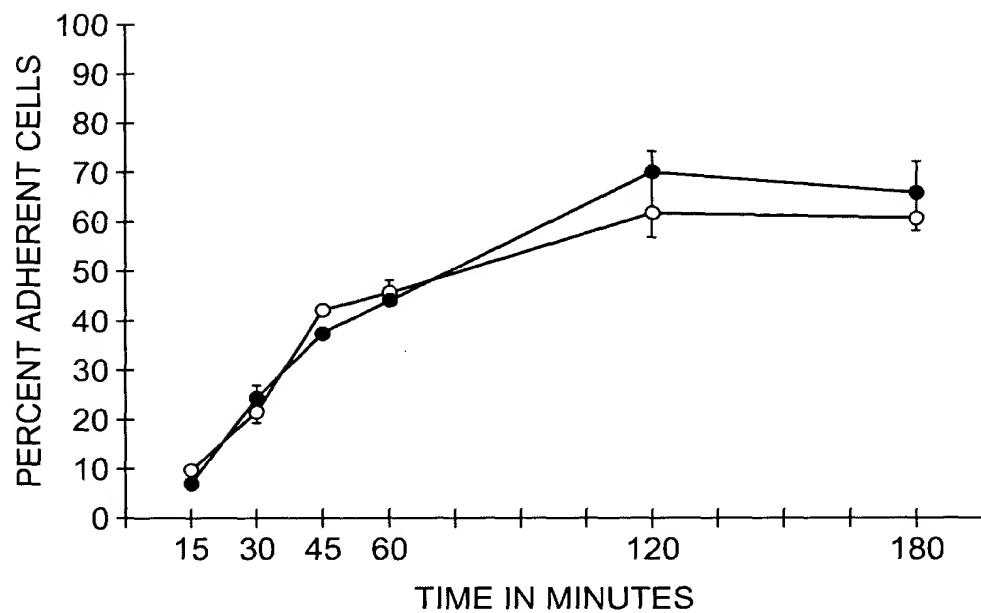
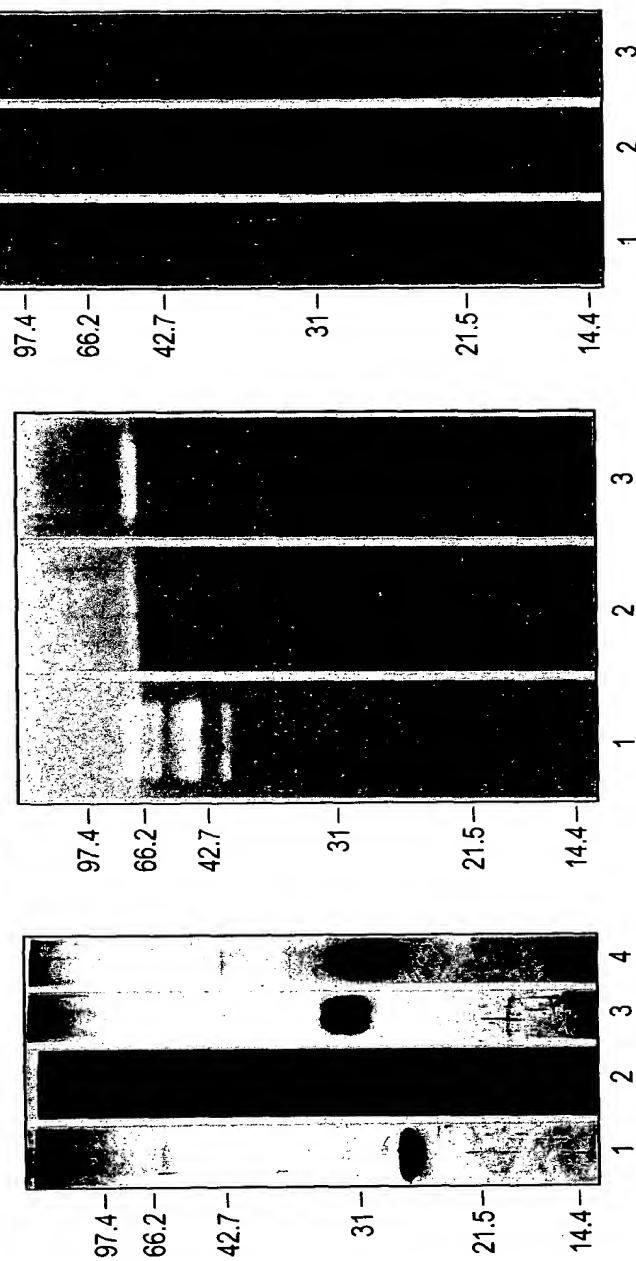


FIG. 18B



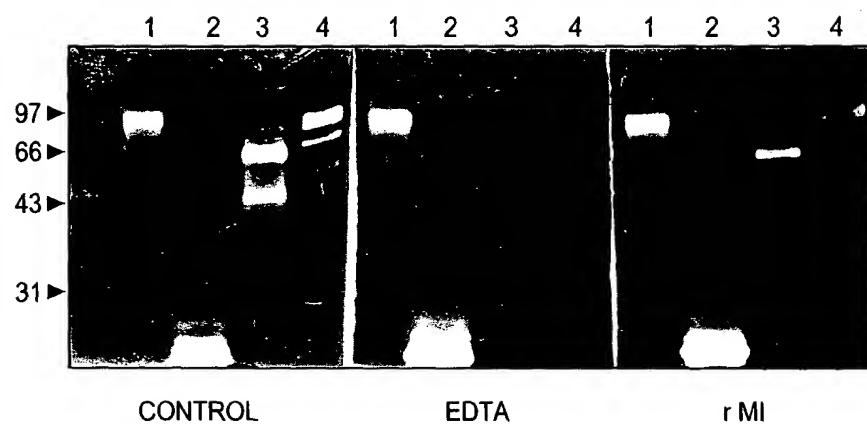
08/26/03

FIG. 5A
FIG. 5B
FIG. 5C



08/26/03

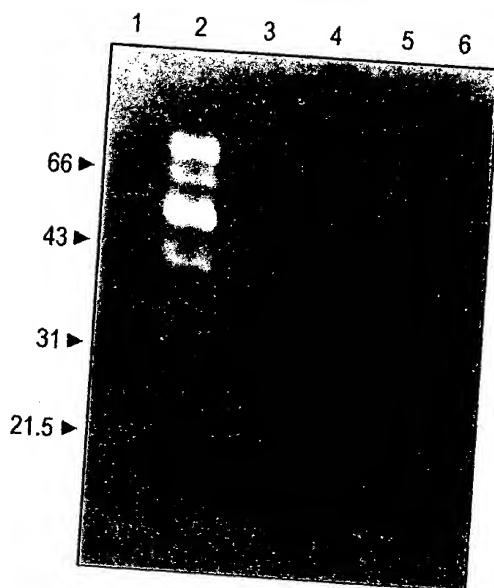
FIG. 14





08/26/03

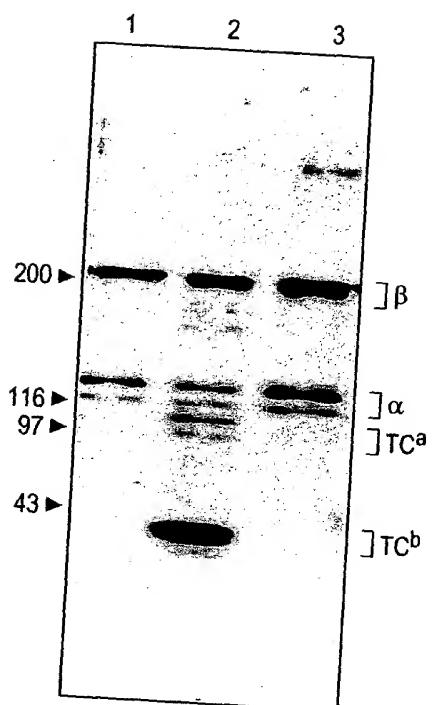
FIG. 13





08/26/03

FIG. 7





08/26/03

FIG. 6

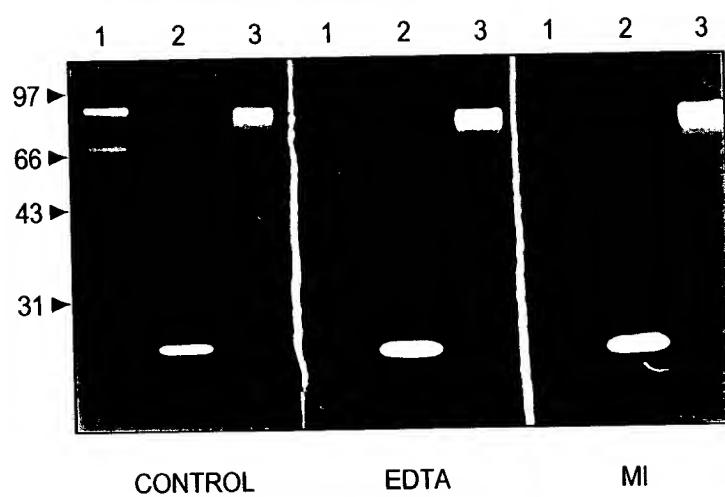


FIG. 19A



FIG. 19B



FIG. 19C

